Release notes for ENDF/B Development n-001_H_001 evaluation



April 26, 2017

• fizcon Warnings:

1. 2-body MT102 OK for 1H: 1H(n,g)2H MAT= 125, MF= 6, MT=102 (1): Is 2-body rxn.

ERROR(S) FOUND IN MAT= 125, MF= 6, MT=102
DISCRETE 2-BODY LAW NOT PERMITTED FOR MT= 102

2. 2-body MT102 OK for 1H: 1H(n,g)2H MAT= 125, MF= 6, MT=102 (2): Is 2-body rxn.

ERROR(S) FOUND IN MAT= 125, MF= 6, MT=102 No problems to report

• fudge-4.0 Warnings:

1. Cross sections do not approach saturation of Wick's limit reactionSuite: (Error # 0): Wick's limit

```
WARNING: Wick's limit too low by 1.672% at 950000.0 WARNING: Wick's limit too low by 4.008% at 1000000.0 WARNING: Wick's limit too low by 8.190% at 1100000.0 WARNING: Wick's limit too low by 11.832% at 1200000.0 ... plus 59 more instances of this message
```

2. Found an invalid distribution for this type of reaction reaction label 1: H2 + photon / Product: photon / Distribution: (Error # 0): Invalid dist.

WARNING: uncorrelated is not a valid distribution component for 2-body reaction

3. Cross section does not match sum of linked reaction cross sections $crossSectionSum\ label\ 0:\ total\ (Error\ \#\ 0):\ CS\ Sum.$

WARNING: Cross section does not match sum of linked reaction cross sections! Max diff: 0.59%

4. The ratio of smallest/largest eigenvalue is quite small, possibly leading to numerical instability in downstream codes.

Section 2 (H2 + photon): / Form 'eval': (Error # 0): Condition num.

WARNING: Ratio of smallest/largest eigenvalue (2.359558e-10) is too small

• fudge-4.0 Errors:

1. Recoil distribution type specified, but recoil partner has unsupported distribution reaction label 1: H2 + photon / Product: H2 / Distribution: / angularTwoBody - recoil: (Error # 0): Bad recoil

WARNING: Recoil distribution type specified, but recoil partner has unsupported distribution type!

• njoy2012 Warnings:

- 1. This nuclide has no URR and NJOY is upset about it unresr...calculation of unresolved resonance cross sections (0): No URR
 - ---message from unresr---mat 125 has no resonance parameters copy as is to nout
- 2. Recoil is not given, so one-particle recoil approximation used. heatr...prompt kerma (0): HEATR/hinit (4)
 - ---message from hinit---mf6, mt102 has recoil with no spectrum photon momentum recoil used.
- 3. This nuclide has no URR and NJOY is upset about it purr...probabalistic unresolved calculation (0): No URR
 - ---message from purr---mat 125 has no resonance parameters copy as is to nout
- 4. The evaluation was missing a file 12. This may be OK. Or not. acer...monte carlo neutron and photon data (1): No MF12
 - message from gamsum---file 12 not found.
- 5. Coefficient mismatch of some sort covr...process covariance data (1): COVR/matshd (2)
 - ---message from matshd---processing of mat/mt 125/ 1 vs. mat1/mt1 125/ 1 largest coefficient= 1.00000E+00 at index 539 555
- 6. The number of coefficients was too large in a covariance covr...process covariance data (2): Cov:Too many coeff.
 - ---message from matshd--- 512 coefficients > 1 reset and continue.
- 7. Coefficient mismatch of some sort covr...process covariance data (3): COVR/matshd (2)
 - ---message from matshd---processing of mat/mt 125/ 1 vs. mat1/mt1 125/ 2 largest coefficient= 1.00000E+00 at index 539 555
- 8. The number of coefficients was too large in a covariance covr...process covariance data (4): Cov:Too many coeff.
 - ---message from matshd--- 512 coefficients > 1 reset and continue.
- 9. Coefficient mismatch of some sort covr...process covariance data (5): COVR/matshd (2)
 - ---message from matshd---processing of mat/mt 125/ 2 vs. mat1/mt1 125/ 2 largest coefficient= 1.00000E+00 at index 539 555
- 10. The number of coefficients was too large in a covariance covr...process covariance data (6): Cov:Too many coeff.
 - ---message from matshd--- 512 coefficients > 1 reset and continue.